



SEQUENCE LISTING

<110> E. I. du Pont de Nemours and Company

<120> Membrane-Bound Desaturases

<130> BB1264

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<150> 60/110,784

<151> 1998-December-03

<160> 17

<170> Microsoft Office 97

<210> 1

<211> 1471

<212> DNA

<213> Picramnia pentandra

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<211> 448

<212> PRT

<213> Picramnia pentandra

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 Ile Ser Lys Trp Thr Lys Glu His Pro Gly Gly Glu Leu Pro Leu Leu
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 Ser Phe Ala Gly Gln Asp Val Thr Asp Ala Phe Ile Ala Tyr His Pro
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 Gly Thr Ala Trp Gln Tyr Leu Asp Arg Phe Phe Thr Gly Tyr Tyr Val
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 Gln Asp Tyr Ser Val Ser Glu Met Ser Lys Asp Tyr Arg Arg Leu Val
 85 90 95
 Ser Glu Phe Ser Lys Met Gly Leu Phe Lys Thr Pro Gly Lys Gly Val
 100 105 110
 Tyr Cys Ser Ile Phe Phe Val Ser Val Leu Phe Ala Leu Ser Val Tyr
 115 120 125
 Gly Val Leu Tyr Cys Lys Ser Thr Trp Ala His Leu Cys Ser Gly Leu
 130 135 140
 Leu Met Gly Met Leu Trp Leu Gln Ser Gly Trp Val Gly His Asp Ser
 145 150 155 160
 Cys His Tyr Gln Val Met Pro Asn Arg Lys Leu Asn Arg Leu Phe Gln
 165 170 175
 Ile Ile Ala Gly Asn Val Ile Ala Gly Val Ser Val Ala Trp Trp Lys
 180 185 190
 Leu Asp His Asn Thr His His Phe Ala Cys Asn Ser Ala Asn Leu Asp
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 Pro Asp Ile Gln His Leu Pro Ile Ile Ala Ile Ser Pro Lys Phe Phe
 210 215 220
 Asn Ser Leu Thr Ser Tyr Tyr His Asn Cys Lys Met Thr Tyr Asp Arg
 225 230 235 240
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 245 250 255
 Leu Leu Ser Val Arg Leu Tyr Leu Phe Ile Leu Ser Phe Lys Val Val
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 Phe Ser Asn Asn Lys Arg Val Tyr Lys Arg Ser Gln Glu Ile Leu Gly
 275 280 285
 Tyr Ala Ala Phe Leu Thr Trp Tyr Ser Leu Leu Ser Arg Leu Pro
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 Asn Trp Pro Glu Arg Val Met Tyr Phe Thr Ser Cys Leu Ala Val Ala
 305 310 315 320

Gly Phe Gln His Trp Gln Phe Ser Leu Asn His Phe Ala Ser Asn Val
325 330 335

Tyr Thr Gly Leu Pro Ser Gly Asn Asp Trp Phe His Gln Gln Thr Lys
340 345 350

Gly Thr Leu Asn Ile Thr Ala Ser Ala Trp Trp Asp Trp Phe His Gly
355 360 365

Gly Leu His Phe Gln Ile Glu His His Leu Phe Pro Arg Met Pro Lys
370 375 380

Cys His Phe Arg Lys Ile Ser Pro Ile Val Asn Lys Leu Cys Gln Lys
385 390 395 400

His Asn Leu Ser Tyr Glu Thr Ala Thr Met Trp Glu Ala Asn Lys Met
405 410 415

Val Tyr Ser Thr Leu Arg Ala Val Ala Met Glu Ala Lys Asp Val Thr
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<211> 1764

<212> DNA

<213> Zea mays

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 <212> PRT
 <213> Zea mays

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Ser Ala Asp Asp Leu Trp Ile Ser Ile Ser Gly Asp Val Tyr Asp Val
 35 40 45

Thr Pro Trp Leu Pro His His Pro Gly Gly Asp Leu Pro Leu Leu Thr
 50 55 60

Leu Ala Gly Gln Asp Ala Thr Asp Ala Phe Ala Ala Tyr His Pro Pro
 65 70 75 80

Ser Ala Arg Pro Leu Leu Arg Arg Phe Phe Val Gly Arg Leu Ser Asp
 85 90 95

Tyr Ala Val Ser Pro Ala Ser Ala Asp Tyr Arg Arg Leu Leu Ala Gln
 100 105 110

Leu Ser Ser Ala Gly Leu Phe Glu Arg Val Gly Pro Thr Pro Lys Val
 115 120 125

Gln Leu Val Leu Met Ala Val Leu Phe Tyr Ala Ala Leu Tyr Leu Val
 130 135 140

Leu Ala Cys Ala Ser Ala Trp Ala His Leu Leu Ala Gly Gly Leu Ile
 145 150 155 160

Gly Phe Val Trp Ile Gln Ser Gly Trp Met Gly His Asp Ser Gly His
 165 170 175

His Arg Ile Thr Gly His Pro Val Leu Asp Arg Val Val Gln Val Leu
 180 185 190

Ser Gly Asn Cys Leu Thr Gly Leu Ser Ile Ala Trp Trp Lys Cys Asn
 195 200 205

His Asn Thr His His Ile Ala Cys Asn Ser Leu Asp His Asp Pro Asp
 210 215 220

Leu Gln His Met Pro Leu Phe Ala Val Ser Pro Lys Leu Phe Gly Asn
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Ile Trp Ser Tyr Phe Tyr Gln Arg Thr Leu Ala Phe Asp Ala Ala Ser
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Lys Phe Phe Ile Ser Tyr Gln His Trp Thr Phe Tyr Pro Val Met Cys
 260 265 270

Ile Ala Arg Ile Asn Leu Leu Ala Gln Ser Ala Leu Phe Val Leu Thr
275 280 285

Glu Lys Arg Val Pro Gln Arg Leu Leu Glu Ile Ala Gly Val Ala Thr
290 295 300

Phe Trp Ala Trp Tyr Pro Leu Leu Val Ala Ser Leu Pro Asn Trp Trp
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Glu Arg Val Ala Phe Val Leu Phe Ser Phe Thr Ile Cys Gly Ile Gln
325 330 335

His Val Gln Phe Cys Leu Asn His Phe Ser Ser Asp Val Tyr Val Gly
340 345 350

Pro Pro Lys Gly Asn Asp Trp Phe Glu Lys Gln Thr Ala Gly Thr Leu
355 360 365

Asp Ile Leu Cys Ser Pro Trp Met Asp Trp Phe His Gly Gly Leu Gln
370 375 380

Phe Gln Ile Glu His His Leu Phe Pro Arg Leu Pro Arg Cys His Leu
385 390 395 400

Arg Lys Val Ala Pro Ala Val Arg Asp Leu Cys Lys Lys His Gly Leu
405 410 415

Thr Tyr Ser Ala Ala Thr Phe Trp Gly Ala Asn Val Leu Thr Trp Lys
420 425 430

Thr Leu Arg Ala Ala Leu Gln Ala Arg Thr Ala Thr Ser Gly Gly
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<212> DNA
<213> Glycine max

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<212> PRT
<213> Glycine max

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Ala Ala Ser Ser Asp Tyr Arg Lys Leu Phe Ser Asp Leu Ser Ala Leu
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Asn Leu Phe Asn Arg Lys Gly His Thr Thr Ser Ile Leu Leu Ser Leu
35 40 45

Ile Leu Thr Leu Phe Pro Leu Ser Val Cys Gly Val Leu Phe Ser Asp
50 55 60

Ser Thr Phe Val His Val Leu Ser Ala Ala Leu Ile Gly Phe Leu Trp
65 70 75 80

Ile Gln Ser Gly Trp Ile Gly His Asp Ser Gly His Tyr Asn Val Met
85 90 95

Leu Ser Arg Arg Leu Asn Arg Ala Ile Gln Ile Leu Ser Gly Asn Ile
100 105 110

Leu Ala Gly Ile Ser Ile Gly Trp Trp Lys Trp Asn His Asn Ala His
115 120 125

His Ile Ala Cys Asn Ser Leu Asp Tyr Asp Pro Asp Leu Gln His Met
130 135 140

Pro	Val	Phe	Ala	Val	Ser	Ser	Arg	Phe	Phe	Asn	Ser	Ile	Thr	Ser	His
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Xaa Tyr Gly Arg Lys Xaa Glu Phe Asp Xaa Ile Ala Xaa Phe Leu Ile
 165 170 175

Cys Tyr Gln His Phe Thr Phe Tyr Pro Val Met Cys Val Ala Arg Val
180 185 190

Asn Leu Tyr Leu Gln Thr Ile Leu Leu Leu Phe Ser Arg Xaa Lys Val
195 200 205

Gln Asp Arg Ala Leu Asn Ile Met Gly Ile Leu Val Phe Trp Thr Trp
210 215 220

Phe Leu Phe Leu Leu Ala Leu Leu Phe Val Pro Ile Gln His Ile Gln
 225 230 235 240

Phe Trp Leu Asn His Leu Ala Glu Asn Leu Tyr Xaa Gly
245 250

<210> 7
<211> 1934
<212> DNA
<213> Glycine max

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<211> 450
<212> PRT
<213> Glycine max

<400> 8
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Val Tyr Asn Val Ser Asp Trp Val Lys Glu His Pro Gly Gly Asp Val
35 40 45

Pro Ile Ser Asn Leu Ala Gly Gln Asp Val Thr Asp Ala Phe Ile Ala
50 55 60

Tyr His Pro Gly Thr Ala Trp Ser His Leu Glu Lys Phe Phe Thr Gly
65 70 75 80

Tyr His Leu Ser Asp Phe Lys Val Ser Glu Val Ser Lys Asp Tyr Arg
85 90 95

Lys Leu Ala Ser Glu Phe Ser Lys Leu Gly Leu Phe Asp Thr Lys Gly
100 105 110

His Val Thr Ser Cys Thr Leu Ala Ser Val Ala Val Met Phe Leu Ile
115 120 125

Val Leu Tyr Gly Val Leu Arg Cys Thr Ser Val Trp Ala His Leu Gly
130 135 140

Ser Gly Met Leu Leu Gly Leu Leu Trp Met Gln Ser Ala Tyr Val Gly
145 150 155 160

His Asp Ser Gly His Tyr Val Val Met Thr Thr Asn Gly Phe Asn Lys
165 170 175

Val Ala Gln Ile Leu Ser Gly Asn Cys Leu Thr Gly Ile Ser Ile Ala
180 185 190

Trp Trp Lys Trp Thr His Asn Ala His His Ile Ala Cys Asn Ser Leu
195 200 205

Asp His Asp Pro Asp Leu Gln His Met Pro Val Phe Ala Val Ser Ser
210 215 220

Arg Phe Phe Asn Ser Ile Thr Ser His Phe Tyr Gly Arg Lys Leu Glu
225 230 235 240

Phe Asp Phe Ile Ala Arg Phe Leu Ile Cys Tyr Gln His Phe Thr Phe
245 250 255

Tyr Pro Val Met Cys Val Ala Arg Val Asn Leu Tyr Leu Gln Thr Ile
260 265 270

Leu Leu Leu Phe Ser Arg Arg Lys Val Gln Asp Arg Ala Leu Asn Ile
275 280 285

Met Gly Ile Leu Val Phe Trp Thr Trp Phe Pro Leu Leu Val Ser Cys
` 290 295 300

Leu Pro Asn Trp Pro Glu Arg Val Met Phe Val Leu Ala Ser Phe Ala
305 310 315 320

Val Cys Ser Ile Gln His Ile Gln Phe Cys Leu Asn His Phe Ala Ala
325 330 335

Asn Val Tyr Val Gly Pro Pro Ser Gly Asn Asp Trp Phe Glu Lys Gln
340 345 350

Thr Ser Gly Thr Leu Asp Ile Ser Cys Ala Ser Ser Met Asp Trp Phe
355 360 365

Phe Gly Gly Leu Gln Phe Gln Leu Glu His His Leu Phe Pro Arg Leu
370 375 380

Pro Arg Cys Gln Leu Arg Lys Ile Ser Pro Leu Val Ser Asp Leu Cys
385 390 395 400

Lys Lys His Asn Leu Pro Tyr Arg Ser Leu Ser Phe Trp Glu Ala Asn
405 410 415

Gln Trp Thr Ile Arg Thr Leu Arg Thr Ala Ala Leu Gln Ala Arg Asp
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Leu Thr Asn Pro Ala Pro Lys Asn Leu Leu Trp Glu Ala Val Asn Thr
435 440 445

His Gly
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<210> 9

<211> 1972

<212> DNA

<213> Triticum aestivum

<400> 9

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ccaccaaagg ggaacgactg gttttagagg caaacagcgg gcacacttgc tatcaagtgc 1260
tccccgtgga tggattgggtt ccatgggtgt ctgcagttcc aggttgaaca ccatttgggtt 1320
cctcgccctgc ctcgctgcca ctataggatg gtcgcgcggta ttgtgcgtga cctttgcaga 1380
aagcatgggc tgctttatgg tgccgcaca ttctgggagg caaatgtaat gacatggaa 1440
acgctaaggg ctgcagcatt gcaggccagg gaagccacta ctggagctgc tccaaagaat 1500
ctggctctgg aagctttgaa cactcatgga tgactggat caggactgga gtatgagaca 1560
attgtaaagcg tcgagccttg cgtgcattgc gttatctgtat tgcttctcgat ttgcgttagag 1620
atattgatcc ttttagctgt tggaaatcgat ttggatttt cgtgttgcga ggtgactatc 1680
tttgcagttc aatcgtgggt tcattgcattca gttgtgtact tgtacaccat atttagattg 1740
ttgggttctc cctatcatgg taactacatc aatactactt gatttacatc ataaaatccg 1800
tggcttatct ttacatccat ttcathttgc ttgcaagttc atgaaactgt aaactcaatt 1860
gatggtttgt agcgtgtata tcctgcgtct atggcagctt gaactgcatt ttggaaacat 1920
gacgattcca ataataaaacg tttagacatt ttctaaaaaaaaaa aaaaaaaaaaa aa 1972

<210> 10
<211> 469
<212> PRT
<213> Triticum aestivum

<400> 10
Met Ala Arg Thr Gly Leu Ala Asp Ala Thr Ala Pro Glu Ala Asp Ala
1 5 10 15

Met Pro Ala Ala Ser Lys Asp Ala Ala Asp Val Arg Met Ile Ser Thr
20 25 30

Lys Glu Leu Gln Ala His Ala Ala Asp Asp Leu Trp Ile Ser Ile
35 40 45

Ser Gly Asp Val Tyr Asp Val Thr Pro Trp Leu Arg His His Pro Gly
50 55 60

Gly Glu Val Pro Leu Ile Thr Leu Ala Gly Gln Asp Ala Thr Asp Ala
65 70 75 80

Phe Met Ala Tyr His Pro Pro Ser Val Arg Pro Leu Leu Arg Arg Phe
85 90 95

Phe Val Gly Arg Leu Ser Asp Tyr Thr Val Pro Pro Ala Ser Ala Asp
100 105 110

Phe Arg Arg Leu Leu Ala Gln Leu Ser Ser Ala Gly Leu Phe Glu Arg
115 120 125

Val Gly His Thr Pro Lys Phe Leu Leu Val Ala Met Ser Val Leu Phe
130 135 140

Cys Ile Ala Leu Tyr Cys Val Leu Ala Cys Ser Ser Thr Gly Ala His
145 150 155 160

Met Phe Ala Gly Gly Leu Ile Gly Phe Ile Trp Ile Gln Ser Gly Trp
165 170 175

Ile Gly His Asp Ser Gly His His Gln Ile Thr Arg His Pro Ala Leu
180 185 190

Asn Arg Leu Leu Gln Val Val Ser Gly Asn Cys Leu Thr Gly Leu Gly
195 200 205

Ile Ala Trp Trp Lys Phe Asn His Asn Thr His His Ile Ser Cys Asn
 210 215 220

Ser Leu Asp His Asp Pro Asp Leu Gln His Leu Pro Leu Phe Ala Val
 225 230 235 240

Ser Thr Lys Leu Phe Asn Asn Leu Trp Ser Val Cys Tyr Glu Arg Thr
 245 250 255

Leu Ala Phe Asp Ala Ile Ser Lys Phe Phe Val Ser Tyr Gln His Trp
 260 265 270

Thr Phe Tyr Pro Val Met Gly Phe Ala Arg Ile Asn Leu Leu Val Gln
 275 280 285

Ser Ile Val Phe Leu Ile Thr Gln Lys Lys Val Arg Gln Arg Trp Leu
 290 295 300

Glu Ile Ala Gly Val Ala Ala Phe Trp Val Trp Tyr Pro Leu Leu Val
 305 310 315 320

Ser Cys Leu Pro Asn Trp Trp Glu Arg Val Ala Phe Val Leu Ala Ser
 325 330 335

Phe Val Ile Thr Gly Ile Gln His Val Gln Phe Cys Leu Asn His Phe
 340 345 350

Ser Ser Ala Val Tyr Val Gly Pro Pro Lys Gly Asn Asp Trp Phe Glu
 355 360 365

Arg Gln Thr Ala Gly Thr Leu Asp Ile Lys Cys Ser Pro Trp Met Asp
 370 375 380

Trp Phe His Gly Gly Leu Gln Phe Gln Val Glu His His Leu Phe Pro
 385 390 395 400

Arg Leu Pro Arg Cys His Tyr Arg Met Val Ala Pro Ile Val Arg Asp
 405 410 415

Leu Cys Lys Lys His Gly Leu Ser Tyr Gly Ala Ala Thr Phe Trp Glu
 420 425 430

Ala Asn Val Met Thr Trp Lys Thr Leu Arg Ala Ala Ala Leu Gln Ala
 435 440 445

Arg Glu Ala Thr Thr Gly Ala Ala Pro Lys Asn Leu Val Trp Glu Ala
 450 455 460

Leu Asn Thr His Gly
 465

<210> 11
 <211> 448
 <212> PRT
 <213> Borago officinalis

<400> 11
 Met Ala Ala Gln Ile Lys Lys Tyr Ile Thr Ser Asp Glu Leu Lys Asn
 1 5 10 15

His	Asp	Lys	Pro	Gly	Asp	Leu	Trp	Ile	Ser	Ile	Gln	Gly	Lys	Ala	Tyr
			20					25					30		
Asp	Val	Ser	Asp	Trp	Val	Lys	Asp	His	Pro	Gly	Gly	Ser	Phe	Pro	Leu
	35					40			45						
Lys	Ser	Leu	Ala	Gly	Gln	Glu	Val	Thr	Asp	Ala	Phe	Val	Ala	Phe	His
	50					55				60					
Pro	Ala	Ser	Thr	Trp	Lys	Asn	Leu	Asp	Lys	Phe	Phe	Thr	Gly	Tyr	Tyr
	65					70			75			80			
Leu	Lys	Asp	Tyr	Ser	Val	Ser	Glu	Val	Ser	Lys	Asp	Tyr	Arg	Lys	Leu
		85					90			95					
Val	Phe	Glu	Phe	Ser	Lys	Met	Gly	Leu	Tyr	Asp	Lys	Lys	Gly	His	Ile
		100					105				110				
Met	Phe	Ala	Thr	Leu	Cys	Phe	Ile	Ala	Met	Leu	Phe	Ala	Met	Ser	Val
		115					120			125					
Tyr	Gly	Val	Leu	Phe	Cys	Glu	Gly	Val	Leu	Val	His	Leu	Phe	Ser	Gly
		130				135				140					
Cys	Leu	Met	Gly	Phe	Leu	Trp	Ile	Gln	Ser	Gly	Trp	Ile	Gly	His	Asp
	145				150				155			160			
Ala	Gly	His	Tyr	Met	Val	Val	Ser	Asp	Ser	Arg	Leu	Asn	Lys	Phe	Met
		165					170			175					
Gly	Ile	Phe	Ala	Ala	Asn	Cys	Leu	Ser	Gly	Ile	Ser	Ile	Gly	Trp	Trp
		180					185			190					
Lys	Trp	Asn	His	Asn	Ala	His	His	Ile	Ala	Cys	Asn	Ser	Leu	Glu	Tyr
		195					200			205					
Asp	Pro	Asp	Leu	Gln	Tyr	Ile	Pro	Phe	Leu	Val	Val	Ser	Ser	Lys	Phe
		210					215			220					
Phe	Gly	Ser	Leu	Thr	Ser	His	Phe	Tyr	Glu	Lys	Arg	Leu	Thr	Phe	Asp
	225				230				235			240			
Ser	Leu	Ser	Arg	Phe	Phe	Val	Ser	Tyr	Gln	His	Trp	Thr	Phe	Tyr	Pro
		245					250				255				
Ile	Met	Cys	Ala	Ala	Arg	Leu	Asn	Met	Tyr	Val	Gln	Ser	Leu	Ile	Met
		260					265			270					
Leu	Leu	Thr	Lys	Arg	Asn	Val	Ser	Tyr	Arg	Ala	His	Glu	Leu	Leu	Gly
		275				280				285					
Cys	Leu	Val	Phe	Ser	Ile	Trp	Tyr	Pro	Leu	Leu	Val	Ser	Cys	Leu	Pro
		290				295				300					
Asn	Trp	Gly	Glu	Arg	Ile	Met	Phe	Val	Ile	Ala	Ser	Leu	Ser	Val	Thr
	305				310				315			320			
Gly	Met	Gln	Gln	Val	Gln	Phe	Ser	Leu	Asn	His	Phe	Ser	Ser	Ser	Val
		325					330			335					

Tyr Val Gly Lys Pro Lys Gly Asn Asn Trp Phe Glu Lys Gln Thr Asp
 340 345 350

Gly Thr Leu Asp Ile Ser Cys Pro Pro Trp Met Asp Trp Phe His Gly
 355 360 365

Gly Leu Gln Phe Gln Ile Glu His His Leu Phe Pro Lys Met Pro Arg
 370 375 380

Cys Asn Leu Arg Lys Ile Ser Pro Tyr Val Ile Glu Leu Cys Lys Lys
 385 390 395 400

His Asn Leu Pro Tyr Asn Tyr Ala Ser Phe Ser Lys Ala Asn Glu Met
 405 410 415

Thr Leu Arg Thr Leu Arg Asn Thr Ala Leu Gln Ala Arg Asp Ile Thr
 420 425 430

Lys Pro Leu Pro Lys Asn Leu Val Trp Glu Ala Leu His Thr His Gly
 435 440 445

<210> 12
 <211> 469
 <212> PRT
 <213> Triticum aestivum

<400> 12
 Met Ala Arg Thr Gly Leu Ala Asp Ala Thr Ala Pro Glu Ala Asp Ala
 1 5 10 15

Met Pro Ala Ala Ser Lys Asp Ala Ala Asp Val Arg Met Ile Ser Thr
 20 25 30

Lys Glu Leu Gln Ala His Ala Ala Asp Asp Leu Trp Ile Ser Ile
 35 40 45

Ser Gly Asp Val Tyr Asp Val Thr Pro Trp Leu Arg His His Pro Gly
 50 55 60

Gly Glu Val Pro Leu Ile Thr Leu Ala Gly Gln Asp Ala Thr Asp Ala
 65 70 75 80

Phe Met Ala Tyr His Pro Pro Ser Val Arg Pro Leu Leu Arg Arg Phe
 85 90 95

Phe Val Gly Arg Leu Thr Asp Tyr Thr Val Pro Pro Ala Ser Ala Asp
 100 105 110

Phe Arg Arg Leu Leu Ala Gln Leu Ser Ser Ala Gly Leu Phe Glu Arg
 115 120 125

Val Gly His Thr Pro Lys Phe Leu Leu Val Ala Met Ser Val Leu Phe
 130 135 140

Cys Ile Ala Leu Tyr Cys Val Leu Ala Cys Ser Ser Thr Gly Ala His
 145 150 155 160

Met Phe Ala Gly Gly Leu Ile Gly Phe Ile Trp Ile Gln Ser Gly Trp
 165 170 175

Ile Gly His Asp Ser Gly His His Gln Ile Thr Arg His Pro Ala Leu
 180 185 190
 Asn Arg Leu Leu Gln Val Val Ser Gly Asn Cys Leu Thr Gly Leu Gly
 195 200 205
 Ile Ala Trp Trp Lys Phe Asn His Asn Thr His His Ile Ser Cys Asn
 210 215 220
 Ser Leu Asp His Asp Pro Asp Leu Gln His Leu Pro Leu Phe Ala Val
 225 230 235 240
 Ser Thr Lys Leu Phe Asn Asn Leu Trp Ser Val Cys Tyr Glu Arg Thr
 245 250 255
 Leu Ala Phe Asp Ala Ile Ser Lys Phe Phe Val Ser Tyr Gln His Trp
 260 265 270
 Thr Phe Tyr Pro Val Met Gly Phe Ala Arg Ile Asn Leu Leu Val Gln
 275 280 285
 Ser Ile Val Phe Leu Ile Thr Gln Lys Lys Val Arg Gln Arg Trp Leu
 290 295 300
 Glu Ile Ala Gly Val Ala Ala Phe Trp Val Trp Tyr Pro Leu Leu Val
 305 310 315 320
 Ser Cys Leu Pro Asn Trp Trp Glu Arg Val Ala Phe Val Leu Ala Ser
 325 330 335
 Phe Val Ile Thr Gly Ile Gln His Val Gln Phe Cys Leu Asn His Phe
 340 345 350
 Ser Ser Ala Val Tyr Val Gly Pro Pro Lys Gly Asn Asp Trp Phe Glu
 355 360 365
 Arg Gln Thr Ala Gly Thr Leu Asp Ile Lys Cys Ser Pro Trp Met Asp
 370 375 380
 Trp Phe His Gly Gly Leu Gln Phe Gln Val Glu His His Leu Phe Pro
 385 390 395 400
 Arg Leu Pro Arg Cys His Tyr Arg Met Val Ala Pro Ile Val Arg Asp
 405 410 415
 Leu Cys Lys Lys His Gly Leu Ser Tyr Gly Ala Ala Thr Phe Trp Glu
 420 425 430
 Ala Asn Val Met Thr Trp Lys Thr Leu Arg Ala Ala Leu Gln Ala
 435 440 445
 Arg Glu Ala Thr Thr Gly Ala Ala Pro Lys Asn Leu Val Trp Glu Ala
 450 455 460
 Leu Asn Thr His Gly
 465
 <210> 13
 <211> 458

<212> PRT

<213> Helianthus annuus

<400> 13

Met Val Ser Pro Ser Ile Glu Val Leu Asn Ser Ile Ala Asp Gly Lys
1 5 10 15

Lys Tyr Ile Thr Ser Lys Glu Leu Lys Lys His Asn Asn Pro Asn Asp
20 25 30

Leu Trp Ile Ser Ile Leu Gly Lys Val Tyr Asn Val Thr Glu Trp Ala
35 40 45

Lys Glu His Pro Gly Gly Asp Ala Pro Leu Ile Asn Leu Ala Gly Gln
50 55 60

Asp Val Thr Asp Ala Phe Ile Ala Phe His Pro Gly Thr Ala Trp Lys
65 70 75 80

His Leu Asp Lys Leu Phe Thr Gly Tyr His Leu Lys Asp Tyr Gln Val
85 90 95

Ser Asp Ile Ser Arg Asp Tyr Arg Lys Leu Ala Ser Glu Phe Ala Lys
100 105 110

Ala Gly Met Phe Glu Lys Lys Gly His Gly Val Ile Tyr Ser Leu Cys
115 120 125

Phe Val Ser Leu Leu Ser Ala Cys Val Tyr Gly Val Leu Tyr Ser
130 135 140

Gly Ser Phe Trp Ile His Met Leu Ser Gly Ala Ile Leu Gly Leu Ala
145 150 155 160

Trp Met Gln Ile Ala Tyr Leu Gly His Asp Ala Gly His Tyr Gln Met
165 170 175

Met Ala Thr Arg Gly Trp Asn Lys Phe Ala Gly Ile Phe Ile Gly Asn
180 185 190

Cys Ile Thr Gly Ile Ser Ile Ala Trp Trp Lys Trp Thr His Asn Ala
195 200 205

His His Ile Ala Cys Asn Ser Leu Asp Tyr Asp Pro Asp Leu Gln His
210 215 220

Leu Pro Met Leu Ala Val Ser Ser Lys Leu Phe Asn Ser Ile Thr Ser
225 230 235 240

Val Phe Tyr Gly Arg Gln Leu Thr Phe Asp Pro Leu Ala Arg Phe Phe
245 250 255

Val Ser Tyr Gln His Tyr Leu Tyr Tyr Pro Ile Met Cys Val Ala Arg
260 265 270

Val Asn Leu Tyr Leu Gln Thr Ile Leu Leu Leu Ile Ser Lys Arg Lys
275 280 285

Ile Pro Asp Arg Gly Leu Asn Ile Leu Gly Thr Leu Ile Phe Trp Thr
290 295 300

Trp Phe Pro Leu Leu Val Ser Arg Leu Pro Asn Trp Pro Glu Arg Val
 305 310 315 320

Ala Phe Val Leu Val Ser Phe Cys Val Thr Gly Ile Gln His Ile Gln
 325 330 335

Phe Thr Leu Asn His Phe Ser Gly Asp Val Tyr Val Gly Pro Pro Lys
 340 345 350

Gly Asp Asn Trp Phe Glu Lys Gln Thr Arg Gly Thr Ile Asp Ile Ala
 355 360 365

Cys Ser Ser Trp Met Asp Trp Phe Phe Gly Gly Leu Gln Phe Gln Leu
 370 375 380

Glu His His Leu Phe Pro Arg Leu Pro Arg Cys His Leu Arg Ser Ile
 385 390 395 400

Ser Pro Ile Cys Arg Glu Leu Cys Lys Lys Tyr Asn Leu Pro Tyr Val
 405 410 415

Ser Leu Ser Phe Tyr Asp Ala Asn Val Thr Thr Leu Lys Thr Leu Arg
 420 425 430

Thr Ala Ala Leu Gln Ala Arg Asp Leu Thr Asn Pro Ala Pro Gln Asn
 435 440 445

Leu Ala Trp Glu Ala Phe Asn Thr His Gly
 450 455

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<210> 14
<211> 38
<212> DNA
<213> Artificial Sequence
  
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<220>
<223> Definition of Artificial Sequence:PCR primer for 5' of pk0011.d5
  
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<400> 14
tttgcggccg caaatcaatg gaagaagcaa agaag 35
  
```

```

<210> 15
<211> 33
<212> DNA
<213> Artificial Sequence
  
```

```

<220>
<223> Definition of Sequence: antisense PCR primer for 3' of pk0011.d5
  
```

```

<400> 15
tttgcggccg ccaggattca cccgaaaatgt ttc 33
  
```

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<210> 16
<211> 823
<212> DNA
<213> Triticum aestivum
  
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<220>
<221> unsure
<222> (48)

<220>
<221> unsure
<222> (538)

<220>
<221> unsure
<222> (686)

<220>
<221> unsure
<222> (704)

<220>
<221> unsure
<222> (717)

<220>
<221> unsure
<222> (727)

<220>
<221> unsure
<222> (729)

<220>
<221> unsure
<222> (737)

<220>
<221> unsure
<222> (741)

<220>
<221> unsure
<222> (750)..(751)

<220>
<221> unsure
<222> (769)

<220>
<221> unsure
<222> (777)

<220>
<221> unsure
<222> (807)

<400> 16
ctcccttaaca aacctccgtt gctgtttaa gatccgatct ccccttcncc cctccccctcc 60
cttcctcctg agtccctgacc accccttcctc gegctccagc taaatccacg ccaccgatgg 120
cccgcacggg ctgcggac gcaacggcgc cggaagccga cgcaatgccg gccgccagca 180
aggacgcgcg cgcgtccgc atgatctcca ccaaggagct gcaggcgcac gcccgcgcgg 240
acgacctctg gatctccatc tccggggacg ttacgacgt cacgccctgg ctgcgccacc 300
acccggggcgg cgaggtcccg ctcatcaccc tcgcggccca ggacgccacc gacgcctca 360
tggcctacca cccggccctcc gtgcggccgc tcctccggcc cttcttcgtc ggccgcctca 420

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ccgactacac tgtccccccc gcctccggc acttccggcg ccttcctcgcg cagctctcct 480
ccgcgggcct ctgcgagcgc gtcggcacac ccccaaggtc ctgtctcgctcg caaagtcnngt 540
gctttctgc atcggcctct actggtcctc gcctgtctaa caccggggcc acatgttcgc 600
cgggggtctca ttgggttatac tggtagtcg ggctggatttgc gcataactccg gcacacaat 660
cacaggcacc tgccctcaacg ctctgnagtg gctcgggaat gctnacggct cggatcnctg 720
gggagtnanc acacaanaca natttgaan ngtcacatac ctgactcana ttccgtntcg 780
ggtcacaagt ctaaaaacttg catcgtnaag acttggtagt cat 823

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<210> 17
<211> 114
<212> PRT
<213> Triticum aestivum

<400> 17
Met Pro Ala Ala Ser Lys Asp Ala Ala Asp Val Arg Met Ile Ser Thr
1 5 10 15

Lys Glu Leu Gln Ala His Ala Ala Asp Asp Leu Trp Ile Ser Ile
 20 25 30

Ser Gly Asp Val Tyr Asp Val Thr Pro Trp Leu Arg His His Pro Gly
35 40 45

Gly Glu Val Pro Leu Ile Thr Leu Ala Gly Gln Asp Ala Thr Asp Ala
50 55 60

Phe	Met	Ala	Tyr	His	Pro	Pro	Ser	Val	Arg	Pro	Leu	Leu	Arg	Arg	Phe	
65					70					75					80	

Phe Val Gly Arg Leu Thr Asp Tyr Thr Val Pro Pro Ala Ser Ala Asp
85 90 95

Phe Arg Arg Leu Leu Ala Gln Leu Ser Ser Ala Gly Leu Phe Glu Arg
100 105 110

Val Gly